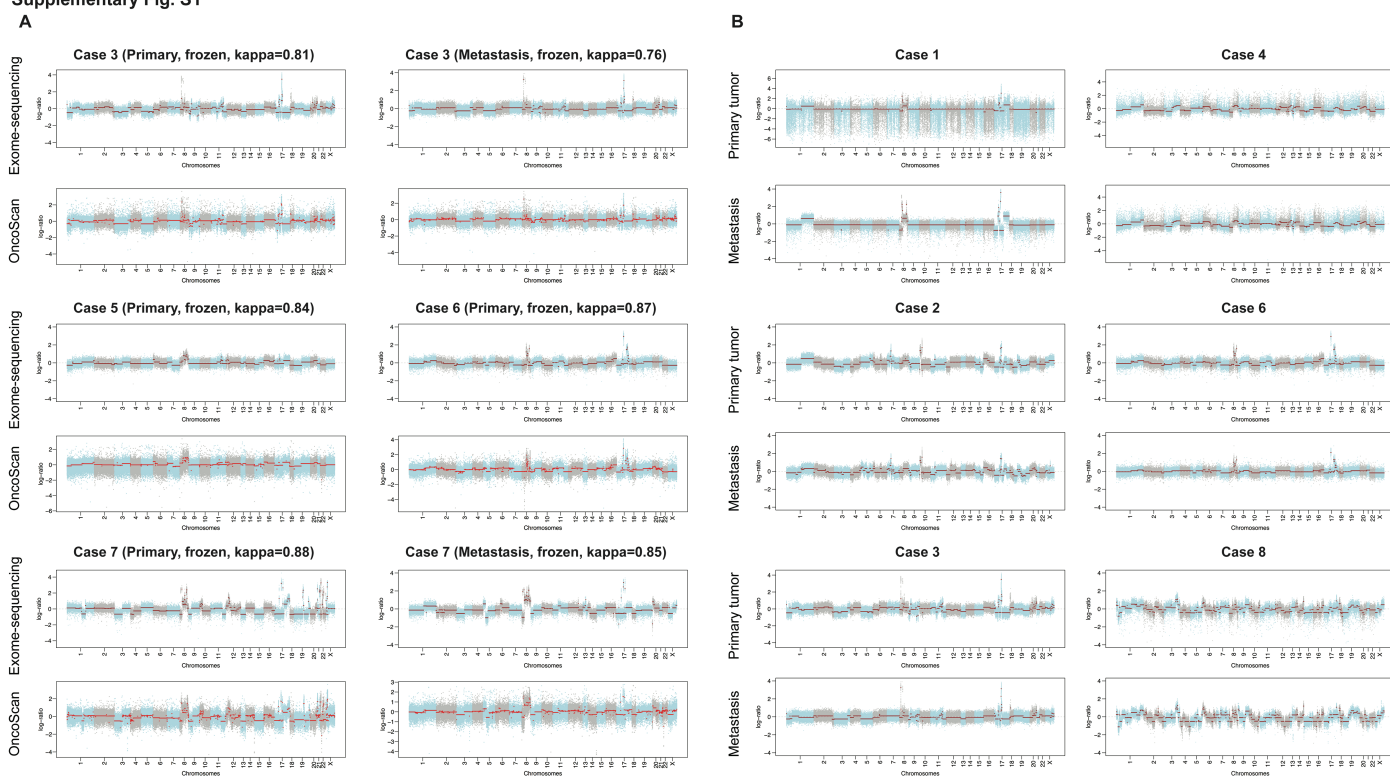


Supplementary Fig. S1



Supplementary Fig. S1: Genome plots of copy number alterations. Genome plots of **(A)** tumors subjected to both whole-exome sequencing and OncoScan for copy number profiling and **(B)** the primary tumors and their synchronous distant metastases for cases 1-4, 6 and 8. In the genome plots, segmented Log_2 ratios (y-axis) were plotted according to their genomic positions (x-axis). Alternating blue and grey demarcate the chromosomes. In **(A)**, agreement of copy number status was assessed using Cohen's weighted kappa statistic.